

SEQUENCE LISTING

<110> Masakazu Takeuchi
Yoshimi Takai

<120> Rabconnectin-3-binding protein

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<150> JP 2002-319521

<151> 2002-11-01

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Ala Pro Thr His Cys Ile Ser Ala Val Leu Leu Thr Asp Asp Gly Ala	
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Thr Ile Val Thr Gly Cys His Asp Gly Gln Ile Cys Leu Trp Asp Leu	
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Ser Val Glu Leu Gln Ile Asn Pro Arg Ala Leu Leu Phe Gly His Thr	
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gca tca atc act tgt ttg tct aaa gct tgt gct tcc agt gac aaa cag	240
Ala Ser Ile Thr Cys Leu Ser Lys Ala Cys Ala Ser Ser Asp Lys Gln	
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Tyr Ile Val Ser Ala Ser Glu Ser Gly Glu Met Cys Leu Trp Asp Val	
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Ser Asp Gly Arg Cys Ile Glu Phe Thr Lys Leu Ala Cys Thr His Thr	
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ggc ata cag ttc tac cag ttc tct gtt ggg aat cag cga gaa gga agg	384
Gly Ile Gln Phe Tyr Gln Phe Ser Val Gly Asn Gln Arg Glu Gly Arg	
115 120 125	
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Leu Leu Cys His Gly His Tyr Pro Glu Ile Leu Val Val Asp Ala Thr	
130 135 140	
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Ser Leu Glu Val Leu Tyr Ser Leu Val Ser Lys Ile Ser Pro Asp Trp	
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Ile Ser Ser Met Ser Ile Ile Arg Ser His Arg Thr Gln Glu Asp Thr	
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Val Val Ala Leu Ser Val Thr Gly Ile Leu Lys Val Trp Ile Val Thr	
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Ser Glu Ile Ser Asp Met Gln Asp Thr Glu Pro Ile Phe Glu Glu Glu	
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Ser Lys Pro Ile Tyr Cys Gln Asn Cys Gln Ser Ile Ser Phe Cys Ala	
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Phe Thr Gln Arg Ser Leu Leu Val Val Cys Ser Lys Tyr Trp Arg Val	
225 230 235 240	
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Phe Asp Ala Gly Asp Tyr Ser Leu Leu Cys Ser Gly Pro Ser Glu Asn	
245 250 255	
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Gly Gln Thr Trp Thr Gly Gly Asp Phe Val Ser Ser Asp Lys Val Ile	
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Ile Trp Thr Glu Asn Gly Gln Ser Tyr Ile Tyr Lys Leu Pro Ala Ser	
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Cys Leu Pro Ala Ser Asp Ser Phe Arg Ser Asp Val Gly Lys Ala Val	
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Glu Asn Leu Ile Pro Pro Val Gln His Ile Leu Leu Asp Arg Lys Asp	
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Arg	Glu	Tyr	Phe	His	Lys	Leu	Leu	Ile	Gln	Gly	Asp	Ser	Ser	Gly	Arg	
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Leu	Asn	Ile	Trp	Asn	Ile	Ser	Asp	Thr	Ala	Asp	Lys	Gln	Gly	Ser	Glu	
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gaa	ggg	ctg	gca	atg	aca	act	tct	att	agt	ttg	caa	gag	gca	ttt	gat	1152
Glu	Gly	Leu	Ala	Met	Thr	Thr	Ser	Ile	Ser	Leu	Gln	Glu	Ala	Phe	Asp	
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Lys	Leu	Asn	Pro	Cys	Pro	Ala	Gly	Ile	Ile	Asp	Gln	Leu	Ser	Val	Ile	
385					390					395					400	
ccc	aat	agt	aat	gaa	cct	ctt	aaa	gta	act	gca	agt	gtg	tac	ata	cca	1248
Pro	Asn	Ser	Asn	Glu	Pro	Leu	Lys	Val	Thr	Ala	Ser	Val	Tyr	Ile	Pro	
				405						410				415		
gca	cat	gga	cga	ctt	gtt	tgt	ggt	cgt	gaa	gat	gga	agc	ata	gtt	att	1296
Ala	His	Gly	Arg	Leu	Val	Cys	Gly	Arg	Glu	Asp	Gly	Ser	Ile	Val	Ile	
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gta	cct	gcc	aca	cag	acg	gcc	ata	gta	cag	ctg	ttg	caa	ggg	gaa	cac	1344
Val	Pro	Ala	Thr	Gln	Thr	Ala	Ile	Val	Gln	Leu	Leu	Gln	Gly	Glu	His	
				435						440				445		
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Met	Leu	Arg	Arg	Gly	Trp	Pro	Pro	His	Arg	Thr	Leu	Arg	Gly	His	Arg	
				450						455				460		
aac	aaa	gtc	aca	tgt	ttg	cta	tat	cct	cat	cag	gtc	tca	gct	cgg	tat	1440
Asn	Lys	Val	Thr	Cys	Leu	Leu	Tyr	Pro	His	Gln	Val	Ser	Ala	Arg	Tyr	
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gat	caa	aga	tac	ctg	ata	tct	gga	ggt	gtg	gat	ttt	tca	gtc	ata	att	1488
Asp	Gln	Arg	Tyr	Leu	Ile	Ser	Gly	Gly	Val	Asp	Phe	Ser	Val	Ile	Ile	
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tgg	gac	ata	ttt	tct	gga	gaa	atg	aaa	cat	atc	ttc	tgt	gtt	cat	ggt	1536
Trp	Asp	Ile	Phe	Ser	Gly	Glu	Met	Lys	His	Ile	Phe	Cys	Val	His	Gly	
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ggt	gag	att	act	caa	ctt	cta	gtt	cca	cct	gaa	aac	tgt	agt	gca	aga	1584
Gly	Glu	Ile	Thr	Gln	Leu	Leu	Val	Pro	Pro	Glu	Asn	Cys	Ser	Ala	Arg	
				515						520				525		
gta	cag	cac	tgc	atc	tgc	tct	gta	gcc	agt	gac	cac	tca	gta	gga	ctt	1632
Val	Gln	His	Cys	Ile	Cys	Ser	Val	Ala	Ser	Asp	His	Ser	Val	Gly	Leu	
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cta	agt	ttg	cga	gag	aaa	aaa	tgc	ata	atg	ttg	gca	tct	cgt	cac	ctt	1680
Leu	Ser	Leu	Arg	Glu	Lys	Lys	Cys	Ile	Met	Leu	Ala	Ser	Arg	His	Leu	

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Phe Pro Ile Gln Val Ile Lys Trp Arg Pro Ser Asp Asp Tyr Leu Val				
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gtg ggg tgt tca gat ggt tct gtg tac gtc tgg caa atg gat act ggt				1776
Val Gly Cys Ser Asp Gly Ser Val Tyr Val Trp Gly				
	580	585	590	
gca ttg gat cgt tgt gtg atg ggg ata aca gca gtt gag att cta aac				1824
Ala Leu Asp Arg Cys Val Met Gly Ile Thr Ala Val Glu Ile Leu Asn				
	595	600	605	
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Ala Cys Asp Glu Ala Val Pro Ala Ala Val Asp Ser Leu Ser His Pro				
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Ala Val Asn Leu Lys Gln Ala Met Thr Arg Arg Ser Leu Ala Ala Leu				
	625	630	635	640
aaa aat atg gcc cat cat aag cta caa acc ctt gca act aac ctc ttg				1968
Lys Asn Met Ala His His Lys Leu Gln Thr Leu Ala Thr Asn Leu Leu				
	645	650	655	
gct tct gag gca tct gac aag gga aat tta cct aaa tat tct cat aac				2016
Ala Ser Glu Ala Ser Asp Lys Gly Asn Leu Pro Lys Tyr Ser His Asn				
	660	665	670	
tcc ctg atg gtt caa gca ata aag aca aac cta aca gac ccg gac ata				2064
Ser Leu Met Val Gln Ala Ile Lys Thr Asn Leu Thr Asp Pro Asp Ile				
	675	680	685	
cat gtg cta ttc ttt gat gtg gaa gcg ttg att att caa ctc ctg act				2112
His Val Leu Phe Phe Asp Val Glu Ala Leu Ile Ile Gln Leu Leu Thr				
	690	695	700	
gaa gaa gcc tct agg ccg aat act gct ctt att tcc cca gag aat ttg				2160
Glu Glu Ala Ser Arg Pro Asn Thr Ala Leu Ile Ser Pro Glu Asn Leu				
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caa aaa gca tct ggc agt tca gac aaa ggg ggc tct ttt tta act gga				2208
Gln Lys Ala Ser Gly Ser Ser Asp Lys Gly Gly Ser Phe Leu Thr Gly				
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aaa cga gca gca gtt ctc ttc caa caa gtg aaa gaa acg atc aaa gag				2256
Lys Arg Ala Ala Val Leu Phe Gln Gln Val Lys Glu Thr Ile Lys Glu				
	740	745	750	
aac atc aag gaa cac ctc ctt gat gat gaa gag gag gat gag gag ata				2304
Asn Ile Lys Glu His Leu Leu Asp Asp Glu Glu Glu Asp Glu Glu Ile				
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Met Arg Gln Arg Arg Glu Glu Ser Asp Pro Glu Tyr Arg Ser Ser Lys				
	770	775	780	

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Ser Lys Pro Leu Thr Leu Leu Glu Tyr Asn Leu Thr Met Asp Thr Ala	
785 790 795 800	
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Lys Leu Phe Met Ser Cys Leu His Ala Trp Gly Leu Asn Glu Val Leu	
805 810 815	
gat gaa gtt tgc ctg gat cgc ctt gga atg ctg aaa ccc cac tgc acc	2496
Asp Glu Val Cys Leu Asp Arg Leu Gly Met Leu Lys Pro His Cys Thr	
820 825 830	
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Val Ser Phe Gly Leu Leu Ser Arg Gly Gly His Met Ser Leu Met Leu	
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ccg ggt tat aat cag cct gct tgt aaa ctg tca cat ggg aaa aca gaa	2592
Pro Gly Tyr Asn Gln Pro Ala Cys Lys Leu Ser His Gly Lys Thr Glu	
850 855 860	
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Val Gly Arg Lys Leu Pro Ala Ser Glu Gly Val Gly Lys Gly Thr Tyr	
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Gly Val Ser Arg Ala Val Thr Thr Gln His Leu Leu Ser Ile Ile Ser	
885 890 895	
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Leu Ala Asn Thr Leu Met Ser Met Thr Asn Ala Thr Phe Ile Gly Asp	
900 905 910	
cat atg aag aag ggt cct acc agg cca cct aga cca agc acc cca gac	2784
His Met Lys Lys Gly Pro Thr Arg Pro Pro Arg Pro Ser Thr Pro Asp	
915 920 925	
ctt tct aag gca agg ggt tcc cct cca act tcc agt aat att gtg caa	2832
Leu Ser Lys Ala Arg Gly Ser Pro Pro Thr Ser Ser Asn Ile Val Gln	
930 935 940	
gga cag att aaa caa gtt gct gca cct gtc gtt tcc gct cgg tct gat	2880
Gly Gln Ile Lys Gln Val Ala Ala Pro Val Val Ser Ala Arg Ser Asp	
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Ala Asp His Ser Gly Ser Asp Pro Pro Ser Ala Pro Ala Leu His Thr	
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Cys Phe Leu Val Asn Glu Gly Trp Ser Gln Leu Ala Ala Met His Cys	
980 985 990	
gtt atg ctg cca gac cta ctg gga ttg gat aaa ttt agg cct ccc ctt	3024
Val Met Leu Pro Asp Leu Leu Gly Leu Asp Lys Phe Arg Pro Pro Leu	
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Leu Glu	Met Leu	Ala Arg	Arg Trp	Gln Asp	Arg Cys	Leu Glu	Val	
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aga gaa	gcc gca	cag gcc	ctg ctt	ctg gcg	gaa ctg	aga aga	att	3114
Arg Glu	Ala Ala	Gln Ala	Leu Leu	Leu Ala	Glu Leu	Arg Arg	Ile	
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gag cag	gca ggc	agg aag	gaa gcc	att gat	gcc tgg	gct cct	tac	3159
Glu Gln	Ala Gly	Arg Lys	Glu Ala	Ile Asp	Ala Trp	Ala Pro	Tyr	
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tta cct	cag tac	ata gac	cac gtc	ata tca	cct gga	gtc aca	tca	3204
Leu Pro	Gln Tyr	Ile Asp	His Val	Ile Ser	Pro Gly	Val Thr	Ser	
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gaa gcc	gcg cag	act atc	acc acg	gct cct	gat gcc	tca ggg	cct	3249
Glu Ala	Ala Gln	Thr Ile	Thr Thr	Ala Pro	Asp Ala	Ser Gly	Pro	
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gaa gca	aaa gtc	cag gag	gaa gag	cat gac	ctt gtt	gac gat	gac	3294
Glu Ala	Lys Val	Gln Glu	Glu Glu	His Asp	Leu Val	Asp Asp	Asp	
1085			1090		1095			
atc acc	act ggt	tgc tta	tca agt	gtc cca	caa atg	aaa aaa	att	3339
Ile Thr	Thr Gly	Cys Leu	Ser Ser	Val Pro	Gln Met	Lys Lys	Ile	
1100			1105		1110			
tct aca	tct tac	gag gaa	aga cgg	aag caa	gct acc	gct att	gtt	3384
Ser Thr	Ser Tyr	Glu Glu	Arg Arg	Lys Gln	Ala Thr	Ala Ile	Val	
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tta ctt	gga gta	ata gga	gct gaa	ttt ggt	gct gaa	att gaa	cct	3429
Leu Leu	Gly Val	Ile Gly	Ala Glu	Phe Gly	Ala Glu	Ile Glu	Pro	
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cct aaa	cta ttg	acc aga	cct cga	agc tct	agc caa	att cct	gag	3474
Pro Lys	Leu Leu	Thr Arg	Pro Arg	Ser Ser	Ser Gln	Ile Pro	Glu	
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gga ttc	ggg ttg	act agt	ggt gga	tcc aac	tac tcg	ctg gcc	aga	3519
Gly Phe	Gly Leu	Thr Ser	Gly Gly	Ser Asn	Tyr Ser	Leu Ala	Arg	
1160			1165		1170			
cat act	tgc aag	gca ctg	acg ttt	ctt ctg	cta cag	cct cca	agc	3564
His Thr	Cys Lys	Ala Leu	Thr Phe	Leu Leu	Leu Gln	Pro Pro	Ser	
1175			1180		1185			
ccc aaa	ctt cct	cca cac	agc act	atc cga	aga aca	gcc att	gat	3609
Pro Lys	Leu Pro	Pro His	Ser Thr	Ile Arg	Arg Thr	Ala Ile	Asp	
1190			1195		1200			
ctg att	gga cgt	ggg ttc	act gtt	tgg gag	cct tac	atg gat	gtg	3654
Leu Ile	Gly Arg	Gly Phe	Thr Val	Trp Glu	Pro Tyr	Met Asp	Val	
1205			1210		1215			
tcc gct	gtt ctg	atg ggg	ctt ctc	gaa ctt	tgt gcc	gat gcc	gag	3699
Ser Ala	Val Leu	Met Gly	Leu Leu	Glu Leu	Cys Ala	Asp Ala	Glu	

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Lys Gln Leu Ala Asn Ile Thr Met Gly Leu Pro Leu Ser Pro Ala			
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Ala Asp Ser Ala Arg Ser Ala Arg His Ala Leu Ser Leu Ile Ala			
1250	1255	1260	
acc gcc aga cca ccc gcc ttc atc acc acc ata gcc aaa gag gta			3834
Thr Ala Arg Pro Pro Ala Phe Ile Thr Thr Ile Ala Lys Glu Val			
1265	1270	1275	
cac aga cat acg gct ctt gca gca aat acc caa tca cag cag aat			3879
His Arg His Thr Ala Leu Ala Ala Asn Thr Gln Ser Gln Gln Asn			
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Met His Thr Thr Thr Leu Ala Arg Ala Lys Gly Glu Ile Leu Arg			
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Val Ile Glu Ile Leu Ile Glu Lys Met Pro Thr Asp Val Val Asp			
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Ser Leu Val Lys Lys Lys Gly Leu Gln Glu Cys Phe Pro Ala Ile			
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Cys Arg Phe Tyr Met Val Ser Tyr Tyr Glu Arg Asn His Arg Ile			
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Thr Gly Lys Cys Gln Thr Ile His Gly His Lys Gly Pro Ile Thr			
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Ala Val Ala Phe Ala Pro Asp Gly Arg Tyr Leu Ala Thr Tyr Ser			
1400	1405	1410	
aac act gac agc cac att tct ttt tgg cag atg aac acg tca ctg			4284
Asn Thr Asp Ser His Ile Ser Phe Trp Gln Met Asn Thr Ser Leu			
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Leu Gly Ser Ile Gly Met Leu Asn Ser Ala Pro Gln Leu Arg Cys			
1430	1435	1440	

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Ile Lys	Thr Tyr	Gln Val	Pro	Pro Val	Gln Pro	Ala	Ser Pro	Gly	
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Ser His	Asn Ala	Leu Lys	Leu	Ala Arg	Leu Ile	Trp	Thr Ser	Asn	
1460			1465			1470			
cgc aac	gtc atc	ctc atg	gcc	cat gac	ggg aag	gag	cac cgc	ttc	4464
Arg Asn	Val Ile	Leu Met	Ala	His Asp	Gly Lys	Glu	His Arg	Phe	
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Met Val									
1490									

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Ser Val Glu Leu Gln Ile Asn Pro Arg Ala Leu Leu Phe Gly His Thr	
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Ala Ser Ile Thr Cys Leu Ser Lys Ala Cys Ala Ser Ser Asp Lys Gln	
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Tyr Ile Val Ser Ala Ser Glu Ser Gly Glu Met Cys Leu Trp Asp Val	
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Ser Asp Gly Arg Cys Ile Glu Phe Thr Lys Leu Ala Cys Thr His Thr	
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Gly Ile Gln Phe Tyr Gln Phe Ser Val Gly Asn Gln Arg Glu Gly Arg	
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Leu Leu Cys His Gly His Tyr Pro Glu Ile Leu Val Val Asp Ala Thr	
	130 135 140
Ser Leu Glu Val Leu Tyr Ser Leu Val Ser Lys Ile Ser Pro Asp Trp	
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Ile Ser Ser Met Ser Ile Ile Arg Ser His Arg Thr Gln Glu Asp Thr	
	165 170 175
Val Val Ala Leu Ser Val Thr Gly Ile Leu Lys Val Trp Ile Val Thr	
	180 185 190

Ser Glu Ile Ser Asp Met Gln Asp Thr Glu Pro Ile Phe Glu Glu Glu
 195 200 205
 Ser Lys Pro Ile Tyr Cys Gln Asn Cys Gln Ser Ile Ser Phe Cys Ala
 210 215 220
 Phe Thr Gln Arg Ser Leu Leu Val Val Cys Ser Lys Tyr Trp Arg Val
 225 230 235 240
 Phe Asp Ala Gly Asp Tyr Ser Leu Leu Cys Ser Gly Pro Ser Glu Asn
 245 250 255
 Gly Gln Thr Trp Thr Gly Gly Asp Phe Val Ser Ser Asp Lys Val Ile
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 Ile Trp Thr Glu Asn Gly Gln Ser Tyr Ile Tyr Lys Leu Pro Ala Ser
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 Cys Leu Pro Ala Ser Asp Ser Phe Arg Ser Asp Val Gly Lys Ala Val
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 Glu Asn Leu Ile Pro Pro Val Gln His Ile Leu Leu Asp Arg Lys Asp
 305 310 315 320
 Lys Glu Leu Leu Ile Cys Pro Pro Val Thr Arg Phe Phe Tyr Gly Cys
 325 330 335
 Arg Glu Tyr Phe His Lys Leu Leu Ile Gln Gly Asp Ser Ser Gly Arg
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 Glu Gly Leu Ala Met Thr Thr Ser Ile Ser Leu Gln Glu Ala Phe Asp
 370 375 380
 Lys Leu Asn Pro Cys Pro Ala Gly Ile Ile Asp Gln Leu Ser Val Ile
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 Ala His Gly Arg Leu Val Cys Gly Arg Glu Asp Gly Ser Ile Val Ile
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 Asp Gln Arg Tyr Leu Ile Ser Gly Gly Val Asp Phe Ser Val Ile Ile
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 Gly Glu Ile Thr Gln Leu Leu Val Pro Pro Glu Asn Cys Ser Ala Arg
 515 520 525
 Val Gln His Cys Ile Cys Ser Val Ala Ser Asp His Ser Val Gly Leu

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Leu Ser Leu Arg Glu Lys Lys Cys Ile Met	Leu Ala Ser Arg His Leu			
545		550		560
Phe Pro Ile Gln Val Ile Lys Trp Arg Pro	Ser Asp Asp Tyr Leu Val			
	565		570	575
Val Gly Cys Ser Asp Gly Ser Val Tyr Val	Trp Gln Met Asp Thr Gly			
	580		585	590
Ala Leu Asp Arg Cys Val Met Gly Ile Thr	Ala Val Glu Ile Leu Asn			
	595		600	605
Ala Cys Asp Glu Ala Val Pro Ala Ala Val	Asp Ser Leu Ser His Pro			
	610		615	620
Ala Val Asn Leu Lys Gln Ala Met Thr Arg	Arg Ser Leu Ala Ala Leu			
625		630		635
Lys Asn Met Ala His His Lys Leu Gln Thr	Leu Ala Thr Asn Leu Leu			
	645		650	655
Ala Ser Glu Ala Ser Asp Lys Gly Asn Leu	Pro Lys Tyr Ser His Asn			
	660		665	670
Ser Leu Met Val Gln Ala Ile Lys Thr Asn	Leu Thr Asp Pro Asp Ile			
	675		680	685
His Val Leu Phe Phe Asp Val Glu Ala Leu	Ile Ile Gln Leu Leu Thr			
	690		695	700
Glu Glu Ala Ser Arg Pro Asn Thr Ala Leu	Ile Ser Pro Glu Asn Leu			
705		710		715
Gln Lys Ala Ser Gly Ser Ser Asp Lys Gly	Gly Ser Phe Leu Thr Gly			
	725		730	735
Lys Arg Ala Ala Val Leu Phe Gln Gln Val	Lys Glu Thr Ile Lys Glu			
	740		745	750
Asn Ile Lys Glu His Leu Leu Asp Asp Glu	Glu Glu Asp Glu Glu Ile			
	755		760	765
Met Arg Gln Arg Arg Glu Glu Ser Asp Pro	Glu Tyr Arg Ser Ser Lys			
	770		775	780
Ser Lys Pro Leu Thr Leu Leu Glu Tyr Asn	Leu Thr Met Asp Thr Ala			
785		790		795
Lys Leu Phe Met Ser Cys Leu His Ala Trp	Gly Leu Asn Glu Val Leu			
	805		810	815
Asp Glu Val Cys Leu Asp Arg Leu Gly Met	Leu Lys Pro His Cys Thr			
	820		825	830
Val Ser Phe Gly Leu Leu Ser Arg Gly Gly	His Met Ser Leu Met Leu			
	835		840	845
Pro Gly Tyr Asn Gln Pro Ala Cys Lys Leu	Ser His Gly Lys Thr Glu			
	850		855	860
Val Gly Arg Lys Leu Pro Ala Ser Glu Gly	Val Gly Lys Gly Thr Tyr			
865		870		875
				880

Gly	Val	Ser	Arg	Ala	Val	Thr	Thr	Gln	His	Leu	Leu	Ser	Ile	Ile	Ser			
				885					890					895				
Leu	Ala	Asn	Thr	Leu	Met	Ser	Met	Thr	Asn	Ala	Thr	Phe	Ile	Gly	Asp			
			900					905					910					
His	Met	Lys	Lys	Gly	Pro	Thr	Arg	Pro	Pro	Arg	Pro	Ser	Thr	Pro	Asp			
		915					920					925						
Leu	Ser	Lys	Ala	Arg	Gly	Ser	Pro	Pro	Thr	Ser	Ser	Asn	Ile	Val	Gln			
	930					935					940							
Gly	Gln	Ile	Lys	Gln	Val	Ala	Ala	Pro	Val	Val	Ser	Ala	Arg	Ser	Asp			
945				950					955					960				
Ala	Asp	His	Ser	Gly	Ser	Asp	Pro	Pro	Ser	Ala	Pro	Ala	Leu	His	Thr			
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Cys	Phe	Leu	Val	Asn	Glu	Gly	Trp	Ser	Gln	Leu	Ala	Ala	Met	His	Cys			
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Val	Met	Leu	Pro	Asp	Leu	Leu	Gly	Leu	Asp	Lys	Phe	Arg	Pro	Pro	Leu			
	995						1000					1005						
Leu	Glu	Met	Leu	Ala	Arg	Arg	Trp	Gln	Asp	Arg	Cys	Leu	Glu	Val	Arg			
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Glu	Ala	Ala	Gln	Ala	Leu	Leu	Leu	Ala	Glu	Leu	Arg	Arg	Ile	Glu	Gln			
1025				1030					1035					1040				
Ala	Gly	Arg	Lys	Glu	Ala	Ile	Asp	Ala	Trp	Ala	Pro	Tyr	Leu	Pro	Gln			
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Tyr	Ile	Asp	His	Val	Ile	Ser	Pro	Gly	Val	Thr	Ser	Glu	Ala	Ala	Gln			
		1060					1065				1070							
Thr	Ile	Thr	Thr	Ala	Pro	Asp	Ala	Ser	Gly	Pro	Glu	Ala	Lys	Val	Gln			
	1075					1080					1085							
Glu	Glu	Glu	His	Asp	Leu	Val	Asp	Asp	Asp	Ile	Thr	Thr	Gly	Cys	Leu			
	1090				1095				1100									
Ser	Ser	Val	Pro	Gln	Met	Lys	Lys	Ile	Ser	Thr	Ser	Tyr	Glu	Glu	Arg			
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Arg	Lys	Gln	Ala	Thr	Ala	Ile	Val	Leu	Leu	Gly	Val	Ile	Gly	Ala	Glu			
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Phe	Gly	Ala	Glu	Ile	Glu	Pro	Pro	Lys	Leu	Leu	Thr	Arg	Pro	Arg	Ser			
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Tyr	Ser	Leu	Ala	Arg	His	Thr	Cys	Lys	Ala	Leu	Thr	Phe	Leu	Leu	Leu			
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Gln	Pro	Pro	Ser	Pro	Lys	Leu	Pro	Pro	His	Ser	Thr	Ile	Arg	Arg	Thr			
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Ala	Ile	Asp	Leu	Ile	Gly	Arg	Gly	Phe	Thr	Val	Trp	Glu	Pro	Tyr	Met			
			1205					1210				1215						
Asp	Val	Ser	Ala	Val	Leu	Met	Gly	Leu	Leu	Glu	Leu	Cys	Ala	Asp	Ala			

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Glu Lys Gln Leu Ala Asn Ile Thr Met Gly Leu Pro Leu Ser Pro Ala			
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Ala Asp Ser Ala Arg Ser Ala Arg His Ala Leu Ser Leu Ile Ala Thr			
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Ala Arg Pro Pro Ala Phe Ile Thr Thr Ile Ala Lys Glu Val His Arg			
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His Thr Ala Leu Ala Ala Asn Thr Gln Ser Gln Gln Asn Met His Thr			
1285	1290	1295	
Thr Thr Leu Ala Arg Ala Lys Gly Glu Ile Leu Arg Val Ile Glu Ile			
1300	1305	1310	
Leu Ile Glu Lys Met Pro Thr Asp Val Val Asp Leu Leu Val Glu Val			
1315	1320	1325	
Met Asp Ile Ile Met Tyr Cys Leu Glu Gly Ser Leu Val Lys Lys Lys			
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Gly Leu Gln Glu Cys Phe Pro Ala Ile Cys Arg Phe Tyr Met Val Ser			
1345	1350	1355	1360
Tyr Tyr Glu Arg Asn His Arg Ile Ala Val Gly Ala Arg His Gly Ser			
1365	1370	1375	
Val Ala Leu Tyr Asp Ile Arg Thr Gly Lys Cys Gln Thr Ile His Gly			
1380	1385	1390	
His Lys Gly Pro Ile Thr Ala Val Ala Phe Ala Pro Asp Gly Arg Tyr			
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Leu Ala Thr Tyr Ser Asn Thr Asp Ser His Ile Ser Phe Trp Gln Met			
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Asn Thr Ser Leu Leu Gly Ser Ile Gly Met Leu Asn Ser Ala Pro Gln			
1425	1430	1435	1440
Leu Arg Cys Ile Lys Thr Tyr Gln Val Pro Pro Val Gln Pro Ala Ser			
1445	1450	1455	
Pro Gly Ser His Asn Ala Leu Lys Leu Ala Arg Leu Ile Trp Thr Ser			
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Met Val			
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